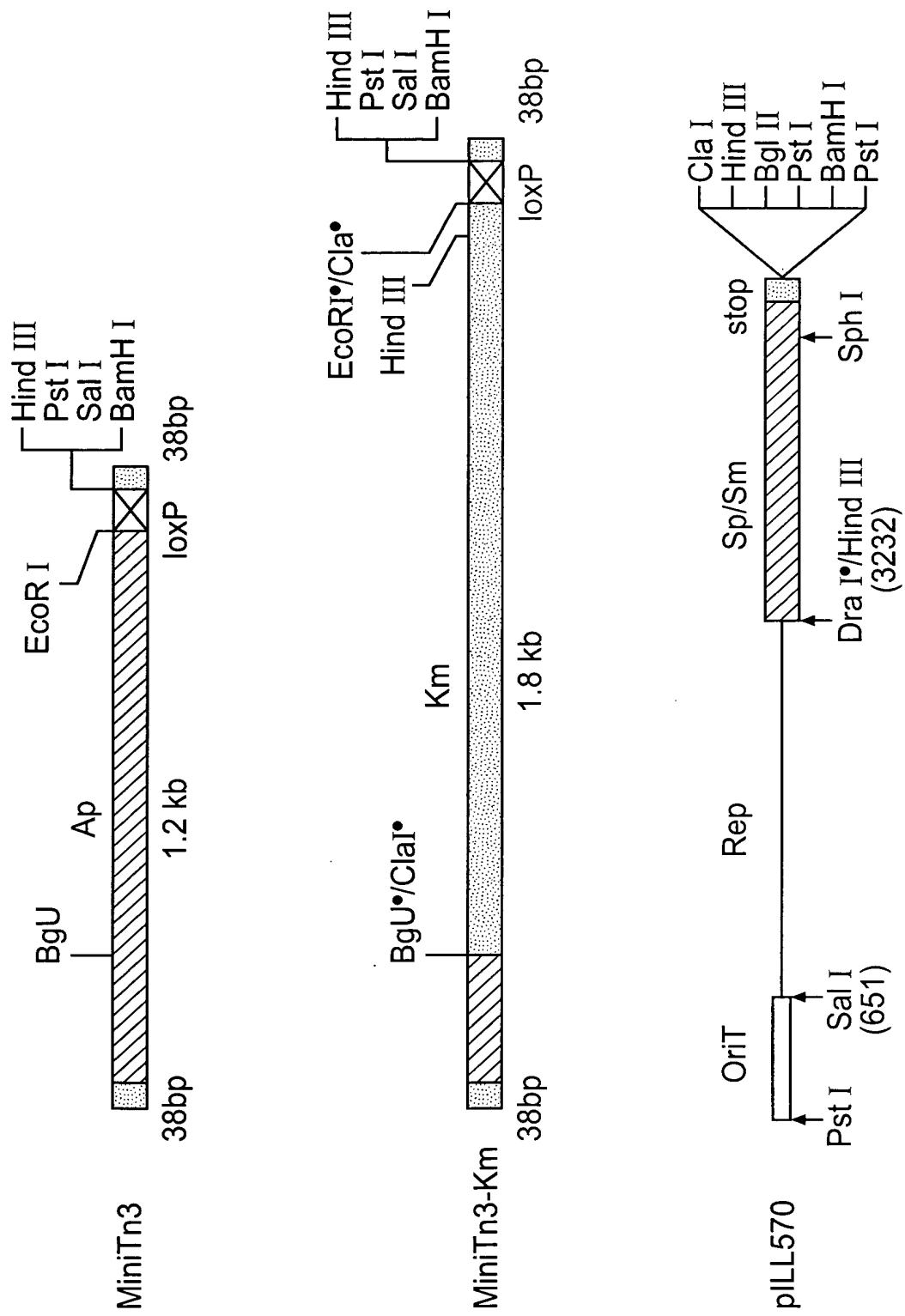


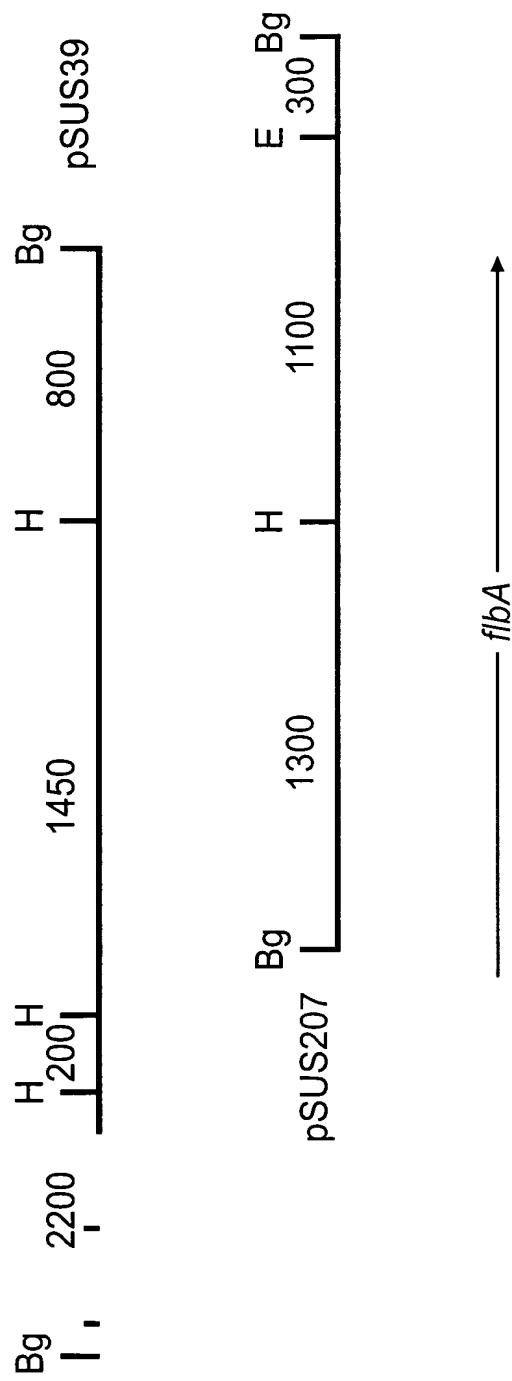


## Replacement Sheet



**FIG. 1A**

Replacement Sheet



**FIG. 1B**



## Replacement Sheet

1	AGC	TTT	GTG	CCA	TAC	TTT	TAA	ACT	TTA	TAT	TAT	AAT	AAG	AGA	CAA	ACA	CAC	CTA	CCA	
61	AAA	TTA	AGG	CAT	TGA	TTT	TAG	ATT	ATG	GCA	AAC	GAA	CGC	TCC	AAA	TTA	GCT	TTT	AAA	AAG
121	ACT	TTC	CCT	GTC	TTT	AAA	CGC	TTC	TTG	CAA	TCC	AAA	GAC	TTA	GCC	CTT	GTG	GTC	TTT	GTG
T	F	P	V	F	K	R	F	L	Q	S	K	D	L	A	L	V	V	F	V	
181	ATA	GCG	ATT	TTA	GCG	ATC	ATT	ATC	GTG	CCG	TTA	CCG	CCT	TTT	GTG	TTG	GAT	TTT	TTA	CTC
I	A	I	L	A	I	I	I	V	P	L	P	P	F	V	L	D	F	L	L	
241	ACG	ATT	TCT	ATC	GCG	CTA	TCG	GTG	TTG	ATT	ATT	TTA	ATC	GGG	CTT	TAT	ATT	GAC	AAA	CCG
T	I	S	I	A	L	S	V	L	I	I	L	I	G	L	Y	I	D	K	P	
301	ACT	GAT	TTT	AGC	GCT	TTC	CCC	ACT	TTA	TTA	CTC	ATT	GTA	ACC	TTA	TAC	CGC	TTG	GCT	TTA
T	D	F	S	A	F	P	T	L	L	L	I	V	T	L	Y	R	L	A	L	
361	AAT	GTC	GCC	ACC	ACT	AGA	ATG	ATT	TTA	ACC	CAA	GGC	TAT	AAA	GGG	CCT	AGC	GCG	GTG	AGC
N	V	A	T	T	R	M	I	L	T	Q	G	Y	K	G	P	S	A	V	S	
421	ATT	ATT	ATC	ACG	GCG	TTT	GGG	GAA	TTT	AGC	GTG	AGC	GGG	AAT	TAT	GTG	ATT	GGG	GCT	ATT
I	I	I	T	A	F	G	E	F	S	V	S	G	N	Y	V	I	G	A	I	

**FIG. 2A**



## Replacement Sheet

481	ATC	TTT	AGT	ATT	TTA	GTG	CTG	GTG	AAT	TTA	TTA	GTG	GTT	ACT	AAT	GGT	TCT	ACT	AGG	GTT
	I	F	S	I	L	V	L	V	N	L	L	V	V	T	N	G	S	T	R	V
541	ACT	GAA	GTT	AGG	GGG	CGA	TTT	GCC	CTA	GAC	GCT	ATG	CCA	GGA	AAG	CAA	ATG	GCG	ATT	GAT
	T	E	V	R	A	R	F	A	L	D	A	M	P	G	K	Q	M	A	I	D
601	GCG	GAT	TTA	AAT	TCA	GGG	CTT	ATT	GAT	GAA	GCT	AAG	GAA	GCT	AAA	AAA	CGG	GCC	GCT	CTA
	A	D	L	N	S	G	L	I	D	K	E	A	K	R	K	R	R	A	A	L
661	AGC	CAA	GAA	GCG	GAT	TTT	TAT	GGT	GGC	ATG	GAT	GGC	GGC	TCT	AAA	TTT	GTC	AAA	GGC	GAT
	S	Q	E	A	D	F	Y	G	A	M	D	G	A	S	K	F	V	K	G	D
721	GCG	ATC	GCT	TCT	ATC	ATT	ATC	ACG	CTT	ATC	AAT	ATC	ATT	GGG	GGT	TTT	TTA	GTG	GGC	GTG
	A	I	A	S	I	I	I	T	L	I	N	I	I	G	G	F	L	V	G	V
781	TTC	CAA	AGG	GAT	ATG	AGC	TTG	AGC	TTT	AGT	GCT	AGC	ACT	TTC	ACT	ATC	TTA	ACC	ATT	GGC
	F	Q	R	D	M	S	L	S	F	S	A	S	T	F	T	I	L	T	I	G
841	GAT	GGG	CTT	GTA	GGG	CAA	ATC	CCT	GCC	TTA	ATC	ATT	GCG	ACA	CGG	ACC	GGT	ATT	GTC	GCC
	D	G	L	V	G	Q	I	P	A	L	I	I	A	T	R	T	G	I	V	A
901	ACT	CGC	ACC	ACG	CAA	AAC	GAA	GAA	GAG	GAC	TTT	GCT	TCT	AAG	CTC	ATC	ACA	CAG	CTC	ACC
	T	R	T	T	Q	N	E	E	D	F	A	S	K	L	I	T	Q	L	T	

**FIG. 2B**



## Replacement Sheet

961	AAT	AAA	AGC	AAA	ACT	TTA	GTG	ATT	GTG	GGG	GCG	ATT	TAT	TGC	TTT	TGC	ACC	ATT	CCT	GGA
N	K	S	K	T	L	V	I	V	G	A	I	Y	C	F	C	T	I	P	G	
1021	CTC	CCT	ACC	TTT	TCT	TTA	GCG	TTT	GTA	GGG	GCT	CTC	TTT	TTA	TTC	ATC	GCA	TGG	CTG	ATT
L	P	T	F	S	L	A	F	V	G	A	L	F	L	F	I	A	W	L	I	
1081	AGC	AGG	GAG	GGA	AAG	GAC	GGG	TTG	CTC	ACT	AAA	TTA	GAA	AAT	TAT	TTG	AGT	CAA	AAA	TTC
S	R	E	G	K	D	G	L	L	T	K	L	E	N	Y	L	S	Q	K	F	
1141	GGC	TTG	GAT	TTG	AGC	GAA	AAA	CCC	CAC	AGC	TCC	AAA	ATC	AAA	CCC	CAC	GCC	CCC	ACC	ACA
G	L	D	L	S	E	K	P	H	S	S	K	I	K	P	H	A	P	T	T	
1201	AGG	GCT	AAA	ACC	CAA	GAA	GAG	ATT	AAA	AGA	GAA	GAG	CAA	GCC	ATT	GAT	GAA	GTG	TTA	
R	A	K	T	Q	E	E	I	K	R	E	E	E	Q	A	I	D	E	V	L	
1261	AAA	ATT	GAA	TTT	TTA	GAA	TTG	GCT	TTA	GGC	TAT	CAG	CTC	TAC	AGC	TTA	GCG	GAC	ATG	AAA
K	I	E	F	L	E	L	A	L	G	Y	Q	L	Y	S	L	A	D	M	K	
1321	CAA	GGG	GGC	GAT	TTG	TTA	GAA	AGG	ATT	AGG	GGT	ATT	AGA	AAA	AAG	ATA	GCG	AGC	GAT	TAT
Q	G	G	G	D	L	L	E	R	I	R	I	G	I	R	K	I	A	S	D	Y
1381	GGT	TTT	TTG	ATG	CCT	CAA	ATT	AGG	ATT	AGG	GAT	AAT	TTA	CAA	CTC	CCC	CCA	ACG	CAT	TAT
G	F	L	M	P	Q	I	R	I	R	D	N	L	Q	L	P	P	T	H	Y	

**FIG. 2C**



## Replacement Sheet

1441	GAA	ATC	AAG	CTT	AAG	GGC	ATT	GTG	ATT	GGT	GAA	GGC	ATG	GTG	ATG	CCG	GAT	AAG	TTT	TTA
E	I	K	L	K	G	I	V	I	G	E	G	M	V	M	P	Q	K	F	L	
1501	GCC	ATG	AAT	ACC	GGT	TTT	GTG	AAT	AAA	GAA	ATT	GAA	GGC	ATT	CCT	ACT	AAA	GAG	CCG	GCT
A	M	N	T	G	F	V	N	K	E	I	E	G	I	P	T	K	E	P	A	
1561	TTT	GGA	ATG	GAC	GCT	TTA	TGG	ATT	GAA	ACT	AAA	AAT	AAA	GAA	GAA	GCC	ATC	ATT	CAA	GGC
F	G	M	D	A	L	W	I	E	T	K	N	K	E	E	A	I	I	Q	G	
1621	TAT	ACC	ATT	ATT	GAT	CCA	AGC	ACC	GTT	ATT	GCG	ACG	CAC	ACC	AGC	GAA	TTA	GTG	AAA	AAA
Y	T	I	I	D	P	S	T	V	I	A	T	H	T	S	E	L	V	K	K	
1681	TAC	GCT	GAA	GAT	TTT	ATC	ACT	AAA	GAT	GAA	GTG	AAA	TCC	CTT	TTA	GAG	CGC	TTG	GCC	AAA
Y	A	E	D	F	I	T	K	D	E	V	K	S	L	L	E	R	L	A	K	
1741/581	GAC	TAT	CCT	ACG	ATT	GTA	GAA	GAG	AGT	AAA	AAA	ATC	CCC	ACC	GGT	GGC	ATC	CGA	TCA	GTC
D	Y	P	T	I	V	E	E	S	K	K	I	P	T	G	A	I	R	S	V	
1801	TTG	CAA	GCC	TTG	TTG	CAT	GAA	AAA	ATC	CCC	ATT	AAA	GAC	ATG	CTC	ACT	ATT	TTA	GAA	ACG
L	Q	A	L	L	H	E	K	I	P	I	K	D	M	L	T	I	L	E	T	
1861	ATT	ACC	GAT	ATT	GGG	CCA	TTA	GTT	CAA	AAC	GAT	GTG	AAT	ATC	TTA	ACC	GAA	CAA	GTG	AGG
I	T	D	I	A	P	L	V	Q	N	D	V	N	I	L	T	E	Q	V	R	
1831	1891																			

**FIG. 2D**



## Replacement Sheet

1921 GCG AGG CTT TCT AGG GTG ATC ACT AAC GCT TTT AAA TCT GAA GAC GGG CGT TTG AAA TTT  
A R L S R V I T N A F K S E D G R L K F  
1981 TTA ACC TTT TCT ACC GAT AGC GAA CAA TTT TTG CTT AAT AAA TTG CGA GAA AAT GGC ACT  
L T F S T D S E Q F L N K L R E N G T  
2041 TCT AAG AGC CTA CTC AAT GTG GGC GAA TTG CAA AAA CTC ATT GAA GCG GTC TCT GAA  
S K S L L N V G E L Q K L I E A V S E  
2101 GAG GCC ATG AAA GTC TTG CAA AAA GGG ATC GCT CCG GTG ATT TTG ATC GTA GAG CCT AAT  
E A M K V L Q K G I A P V I L I V E P N  
2161 TTA AGA AAA GCC CTT TCT AAT CAA ATG GAG CAG ATT GAT GTA ATC GTG CTA AGC  
L R K A L S N Q M E Q A R I D V I V L S  
2221 CAT GCT GAA TTA GAT CCT AAC TCT AAT TTT GAA GCC TTA GGC ACG ATC CAT ATT AAC TTT  
H A E L D P N S N F E A L G T I H I N F  
2281 TAA GGG ATA AAT AAT TGA TAA AAA AGG AGA ATG ATG CAA GTT TAT CAC CTT TCA CAC ATT  
2341 GAT TTA GAC GGC TAT GCA TGC CAG CTT GTT TCA AAA CAA TTT TTT AAA AAT ATC CAA TGC  
2401 TAT AAC GCT AAT TAC GGG CGT GAA GTC TCA GCG AGA ATT TAT GAG ATT TTA AAC GCG ATC  
2461 GCT CAA TCT AAA GAG AGT GAA TTC CTT ATT TTG ATT AGC GA

**FIG. 2E**



## Replacement Sheet

1	MANKRS-KLAFKKKTFPVFKRFLQSKDLALVVFVIAILAIIVPLPPFVLDFLLTISIALS	HpF1bA
1	MAKNKIVDLVFPFLGPLIAPVLAKSLTIVGFLVCILAIIVPLPSPILDFFFLASIALS	CjF1bA
1	MADAAPNASSMPSAKSLLDGLMRGEMGLGVVGIVVLLIIPVPAPLLDVLLAISLTGS	CCF1bF
1	-----MNPHDLEWLNRIGERKDIMLAVLLAVVFMVVPLPVLVLDILIAVNMTIS	YplcrD
1	-----MLLSLLNSARLRRPLLIVLMMITISMFVILPLPTYLVDFLIANNIVLA	StInVA
1	-----MVMIAIMLIIPLPTYLVDFLIGLNVLA	SfMxiA
	.* . * . * . * . * . * . * .	
60	VLIILIGLYIDKPTDFSAFPPTLLLIVTLYRLALNVATTRMILTQGYKGPSAVSIIITAFCG	HpF1bA
61	VLIILISIYIPKPTDLTTFPTLILLITLFRLSLNIAATTRMILSEONGPEAVSEIIIAAFG	CjF1bA
61	VLIILMTAILIKKPLEFTSFPTVLLVTLERLGNIASTRLLILSHGQEGTGGAGAVIEAFG	CCF1bF
52	VLIILMIAIYINSPLQFSAFPAVLLVTLERLALSVSTTRMILLQ-----ADAGQIVVYTFG	YplcrD
49	ILVFMGSFYIDRILSFSTFPAVLLITLFRLALSISTSRLILIEA-----DAGEIIATFG	StInVA
29	ILVFMGSFYIERILSFSTFPVLLITLFRLALSISTSRLILVDAADRGK-----ITTEFG	SfMxiA
	.* . * . * . * . * . * . * . * . * .	
120	EESVSGNYVIGAIIFSIIIVLNLLVVTINGSTRVTEVRAREFALDAMPGKQMAIDADLNSGL	HpF1bA
121	EFVVGGNMVGIVVFCILVLINFMVVTKGSTRVSEVQARFTLDAMPGKQMAIDADLNAGL	CjF1bA
121	HLMMQGNFVIGVIVFIIILIVVNFMVVTKGSGRIAEVAAARFTLDAMPGKQMAIDADLSTGL	CCF1bF
107	NEVVGGNLIVGIVIFIITIVQFLVITKGSERVAEVSAREFSLDAMPGKOMSIDGDMRAGV	YplcrD
104	QFVIGDSSLAVGFVVFVFSIVVVQFIVITKGSERVAEVAAARFSLDGMMPGKQMSIDADLKAGI	StInVA
84	QFVIGDSSLAVGFVVFVFSIVVVQFIVITKGSERVAEVAAARFSLDGMMPGKQMSIDADLKAGI	SfMxiA
	.* . * . * . * . * . * . * . * . * . * .	

**FIG. 3A**



## Replacement Sheet

180 IDDKEAKKRAALSQEADFYGAMDGASKFVKGDAIASIIITLINIIGGFLVGVFQRDMSL HpF1bA  
181 IDEQTARRRQEVTAEANFYGAMDGSSKF1KGDAVAGIIITLINIIGGFLIGSFQHDMAL CjF1bA  
181 ISQDEAKIRRKELEQUESTFGAMDGASKFVKGDAIAGLIITAINIIGGIIIGVVQHKMPF CCF1bF  
167 DVNEARERRATIEKESQMFGSMDGAMKFKVKGDAIAGLIIIFVNILGGVTIGVTQKGAA YplcrD  
164 IDADAARERSVLERESQLYGSFDDGAMKFKIKGDAIAGIIIFVNFIIGGISMVGMTRHGMDL StInVA  
144 IDAAGAKERRSILERESQLYGSFDDGAMKFKIKGDAIAGIIIFVNLIIGGISMQHGMMSL SFMxiA  
\* . \* \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . . . . .  
240 SFSASTFTILTIGDGLVQIPALIIATRTGIVATRTQNEEEDFASKLITQLTNKSKTLLV HpF1bA  
241 SDAASTYTILTIGDGLVSIQPGLITSTATAITITRASKDEENFAEGTLTQILSEYRTLLI CjF1bA  
241 GDAASTYTIMTIGDGLVSIQIPALIIASIAAGMVWSKAGVEGSADKALTQAMNPVGLGMV CCE1bF  
227 AEALQLYSILTVGDMVSVQPVALLIAITAGIIVTRVSSSEDSDLGSDIGKQVVAQPKAMI YplcrD  
224 SSALSTYTMLTIGDGLVAQIPALLIAISAGFIVTRVNGDTDN-MGRNIMTQLNNPFVLV StInVA  
204 SGALSTYTILTIGDGLVSIQIPALLISAGFMLTRVNGDSDN-MGRNIMMSQIFGNPFPVLLI SFMxiA  
\* . . . . . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . \* . . . . .  
300 IVGAIYIC-ECTIPGLPTFSLAFVGALFLFIAWLISREGKDGILLTKLENYLSQFGLDLSE HpF1bA  
301 VGFVLF1-FALVPGLPTLSQLGFMALVFLSLGYLTQVKVEGKI-----DITTVKKSKPSAA CjF1bA  
301 SASSG-I-IALIPGMPIFPFAAMALA-----AGALAY CCE1bF  
287 IGGVLLLFLGLIPGFPTVTFLLIALLVCGGYMLSRKQSRNDEANQDLQSIITSGSGAPA YplcrD  
283 VTAIILTISMGTLPGPFLPVFVILSVVLSVLFYFKFREAKRSAAKPKTSKGEQPLSIEEKE StInVA  
263 VTSALALATIGMILPGFPFFFLIAVTLTALFYKKVVEKEKSLSSESDDSSGGYTG----- SFMxiA  
\* . \* . . . .

**FIG. 3B**



## Replacement Sheet

419 EIRGIRKKIASDYGFLMPQIRIRDNIQLPPTHYEIKLKGIIVIGPGMVMPPDKFLAMNTGF  
 411 EIRSMRRSIAESLGFLMPKIRIRDNLRLKPNYESKLIKGVSIASAEIYPDKYLAAMDGF  
 391 DQIRALRKTLASEYGFVMPVRLDNMRLANQGYAIRIKEMEAGAGEVRLGCLMCMMDPRG  
 392 DELVRVRRALYLDLGVPFPFGIHLRFNEGGMGEGEYIISLQEVPVARGELKAGYLLVRESVS  
 378 QLAERLRSOFFIDYGVRLPEVLLRDGEGLDDNSIVVLLINEIRVEQFTVYFD--LMRVVNY  
 359 GLIERIRSQFFIDYGVRLPTILYRTSNEELKVVDDIVVLLINEVRADSFNIYFDKVCITDENG  
 \* \* \*

479 VNKEIEGIPPTKEPAF--GMDALWIETKNKEEAIQGYTIIDPSTVIATHHTSELVKKYAED HpF1bA  
 471 ITEEIEGIAATKEPAF--NSDALWIDANLKDDEATLNGYIVIDPASVISTHMSELIKAHASE CjF1bA  
 451 GOVELPGEHVREPAF--GLPATWIAADDLREEATFRGYTVIDPATVLTTHILTEILKENMAD CCF1bF  
 452 QLELLGIPYEKGEHLLPDQEAFWVSVVEYERLEKSQLEFFSHSQVLTWHLSHVLRREYAEAD YpLcRD  
 436 SDEWVSSFGINPHTHQOGSSQYFWVTHEEGEKLRELGYVLRNALDELYHCLAVTVARNVNE StInVA  
 419 DIDALGIPVVSTS--YNERVISSWVDVSYTENLTNIDAKIKAQDEFYHQLSQALLNNINE SfMxIA

FIG. 3C



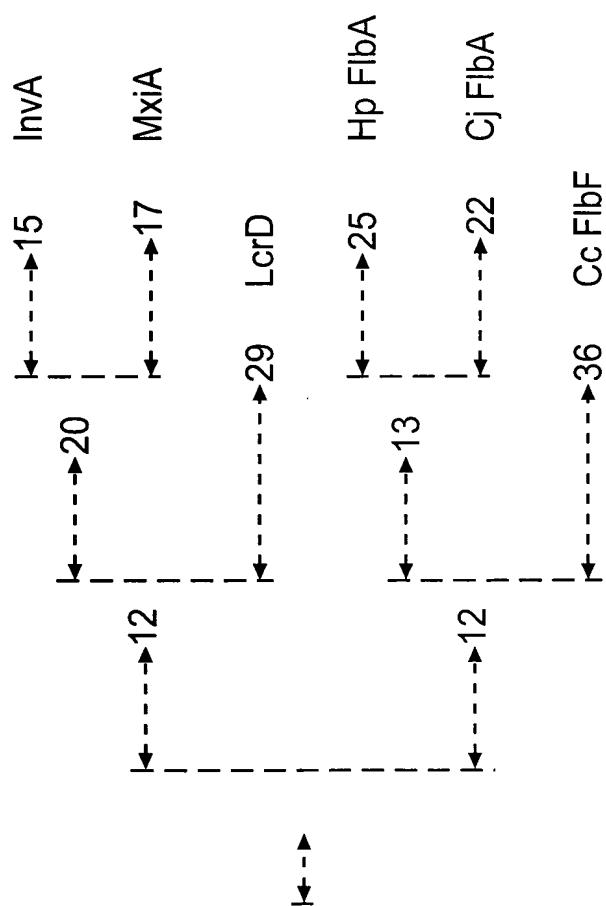
## Replacement Sheet

537	FITKDEVKSSLERLAKDYPPTIVEESK-KI-PTGAI	RSVLOQALLHEKIPIKDMLTILETIT	HpE1bA
529	LITRQEVNLLDKVKNDYPPIVEGAL-GVAPV	SLIQQKILKDLLKHHIPIKDMLTILESVS	CJF1bA
509	LLSYAEVQKLLKEPLPETQKKLVDDLI	PGTVQRLQSLLIREVSVIRLQ	CCF1bF
512	FIGIQETRYLLEQMEGGYGE	LLIREVSVIRLQRLVGEDISIRNMRSILEAMV	YPLCRD
496	YFGIQETKHMDOLEAKFPDLILKEVLRH-	IVPLQRMTEILQRLSERVSVRNMKLIMEALA	STINVA
477	IEGIQETKNMIDQENRYPDLLIKEVFRHV-	IQRISEVLQRLIGENISVRNLKLMESLA	SFMXIA
*	*	*	*
595	DIAPLVONDVNILTEOVRARLSRVITNAPKSE	DGRLKFKELTESTDSEQELLNLBENGTSK	HpE1bA
588	DIAR-VSKSFDMIEKVRASLARMITNMYLDDKG	NLDIFLDSASSAVLMEVNQFRDGSY	CJF1bA
569	EAAPHTA-SVTQLVEQVRARLARQLCWANR	GDDGALPITLSDADWEQAFAEALIGPGDDK	CCF1bF
571	EWGQK-EKDVVQLTETYIRSSILKRYTCY	KYANGNNILPAYLFDOEVEEKIRSGVRQTSAGS	YPLCRD
555	LWAPR-EKDVINLVEHIRGAMARYTICHKE-	ANGGELRAVMVSAEVEDVIRKGIRQTSGSS	STINVA
536	LWAPR-EKDVTITLVEHVRASISRYTICK-	IAVSGETKVVMLSGYIEDAIRKGIRQTSGGS	SFMXIA
*	*	*	*
655	SLLNVGEIQLLIFAVSEAMKVVLQKGIA	PVLKALSNQMEQARIDVVLSHA	HpE1bA
647	HPLLSVAQTGTLVDLRAEVAAVANGRIKPF	ILLCVPQLRKFIADICYNFSINIVVLSFA	CJF1bA
628	QLALPPSRLLQDFIRGVRSFERAALAGEAPV	ILL-TSPGVRPYVRSIIERFRGOTVVMSON	CCF1bF
630	YLALEPAVTESLLEQVRKTIGDILSQIQS	KP-VLIVSMDIRRYVVRKLIIESEYYGLPVL	YPLCRD
613	FLSLDPFASANLMDLITLKLDDLLIAH-	SVDRRRFIKKMIEGRFPDLEVLSEFG	STINVA
594	FLNMDIEVSDEVMETLAHALREL-RNAKK	FLNREKSLDNREKSLDNREKSLDNREKSL	SFMXIA
*	*	*	*
715	ELDPNSNFEALGTHINF		HpE1bA
707	EIAENTNENTEGITIRIEL		CJF1bA
687	EIHPRARLKTVGMV---		CCF1bF
689	ELTQINIQPLGRICL--		YPLCRD
672	EIADSKSVNVIKTI----		STINVA
653	EIDEAYTINVLKTI----		SFMXIA
*	*	*	*

**FIG. 3D**



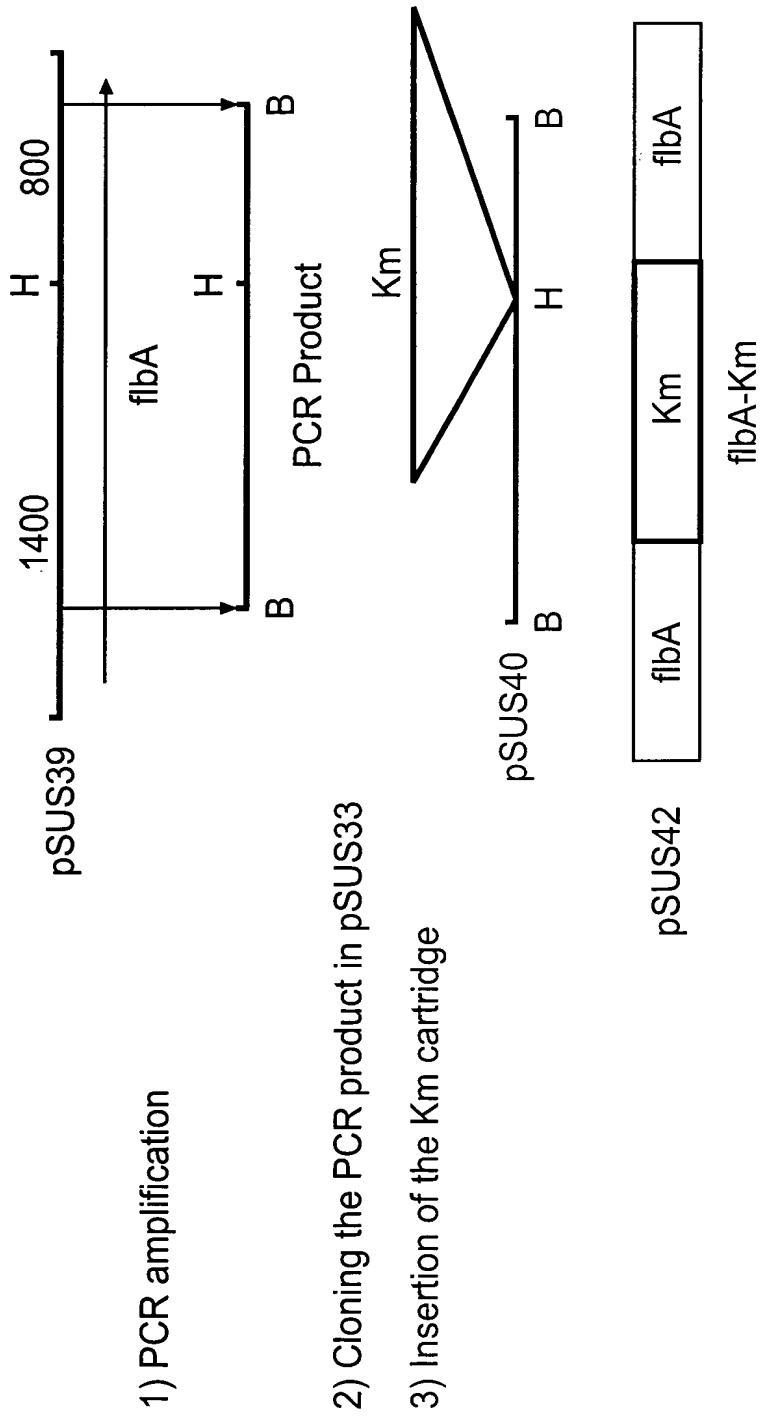
## Replacement Sheet



**FIG. 4**



## Replacement Sheet



**FIG. 5**



# Replacement Sheet

BEST AVAILABLE COPY

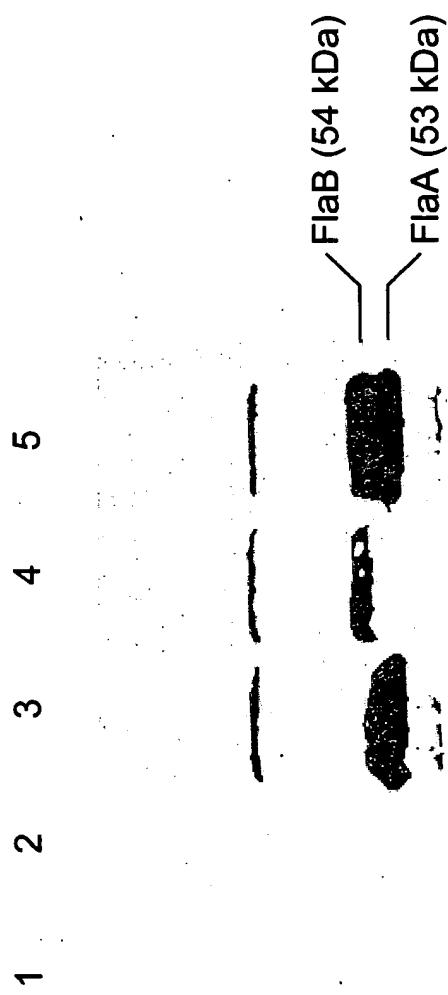
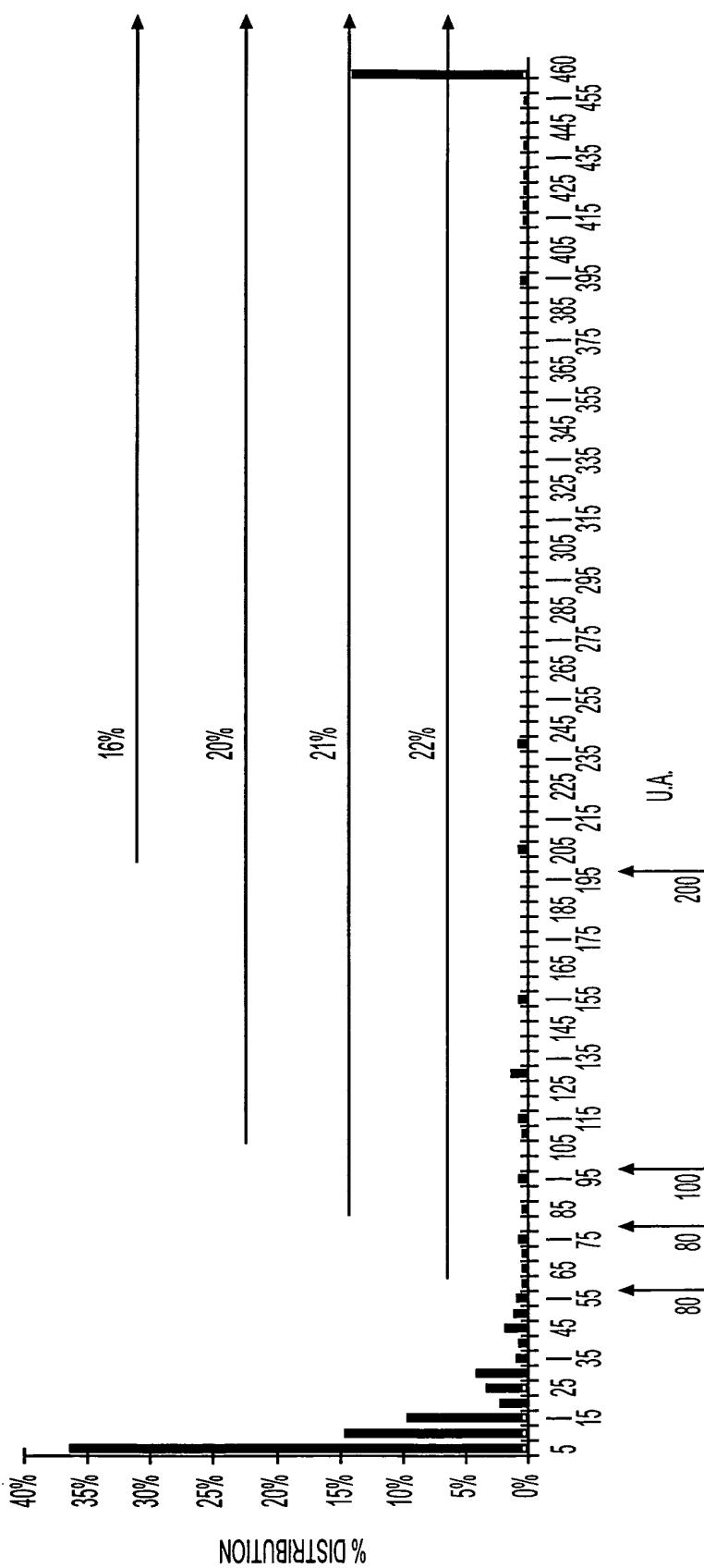


FIG. 6



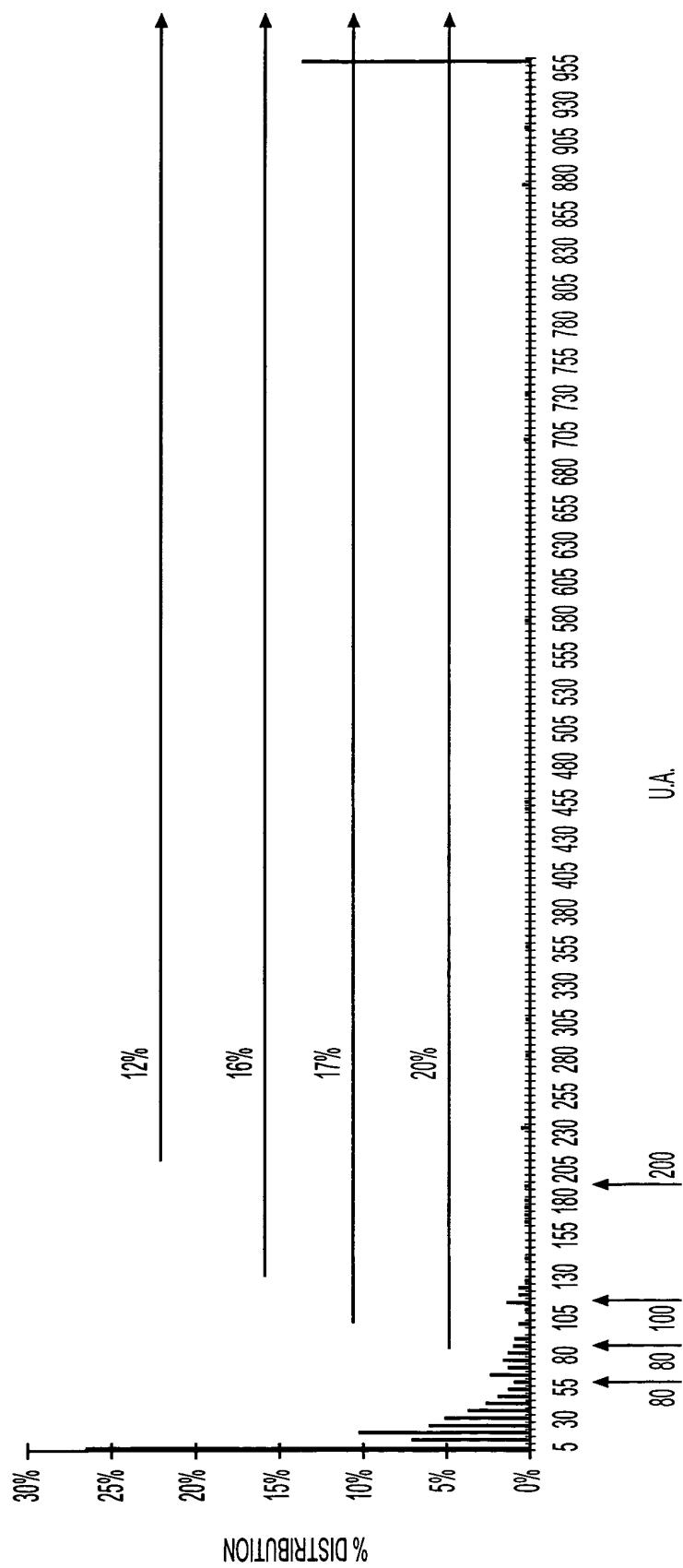
## Replacement Sheet



**FIG. 7**

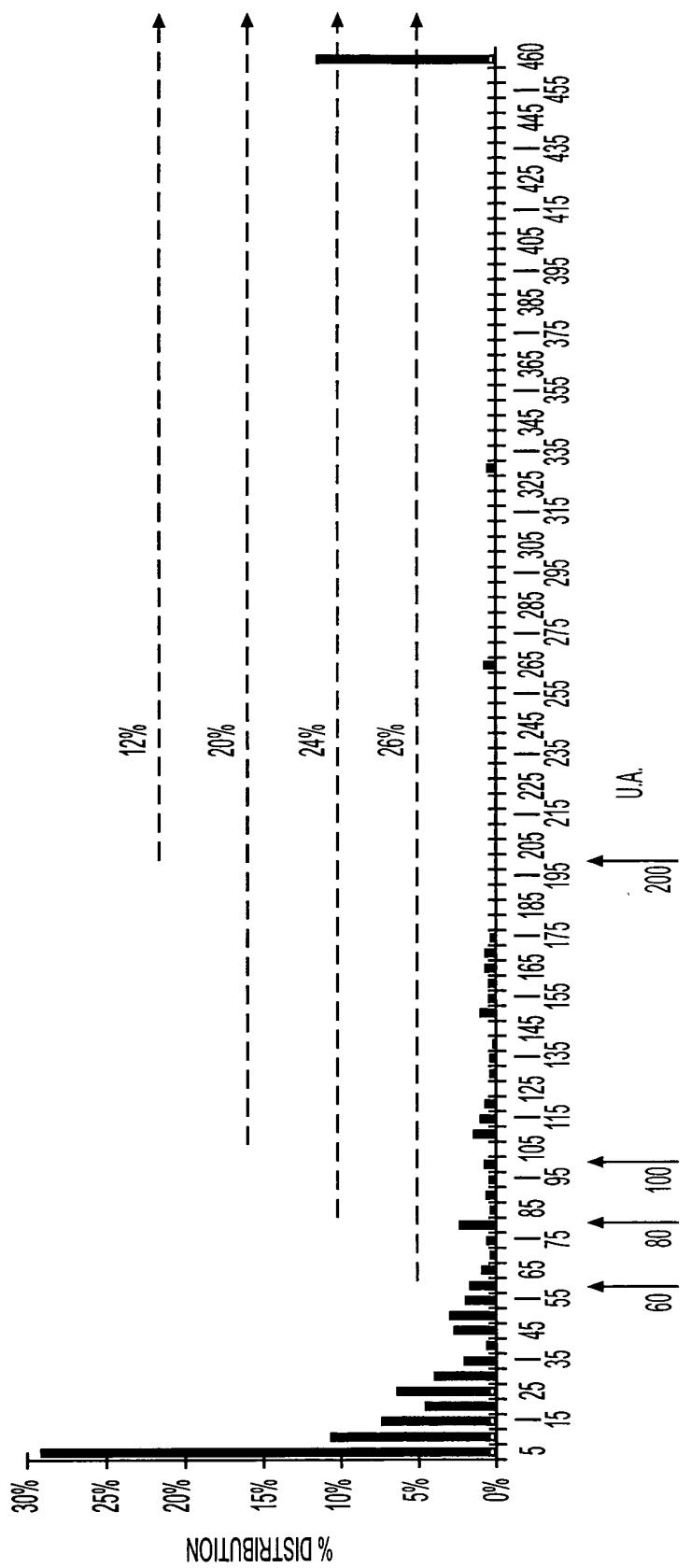


## Replacement Sheet



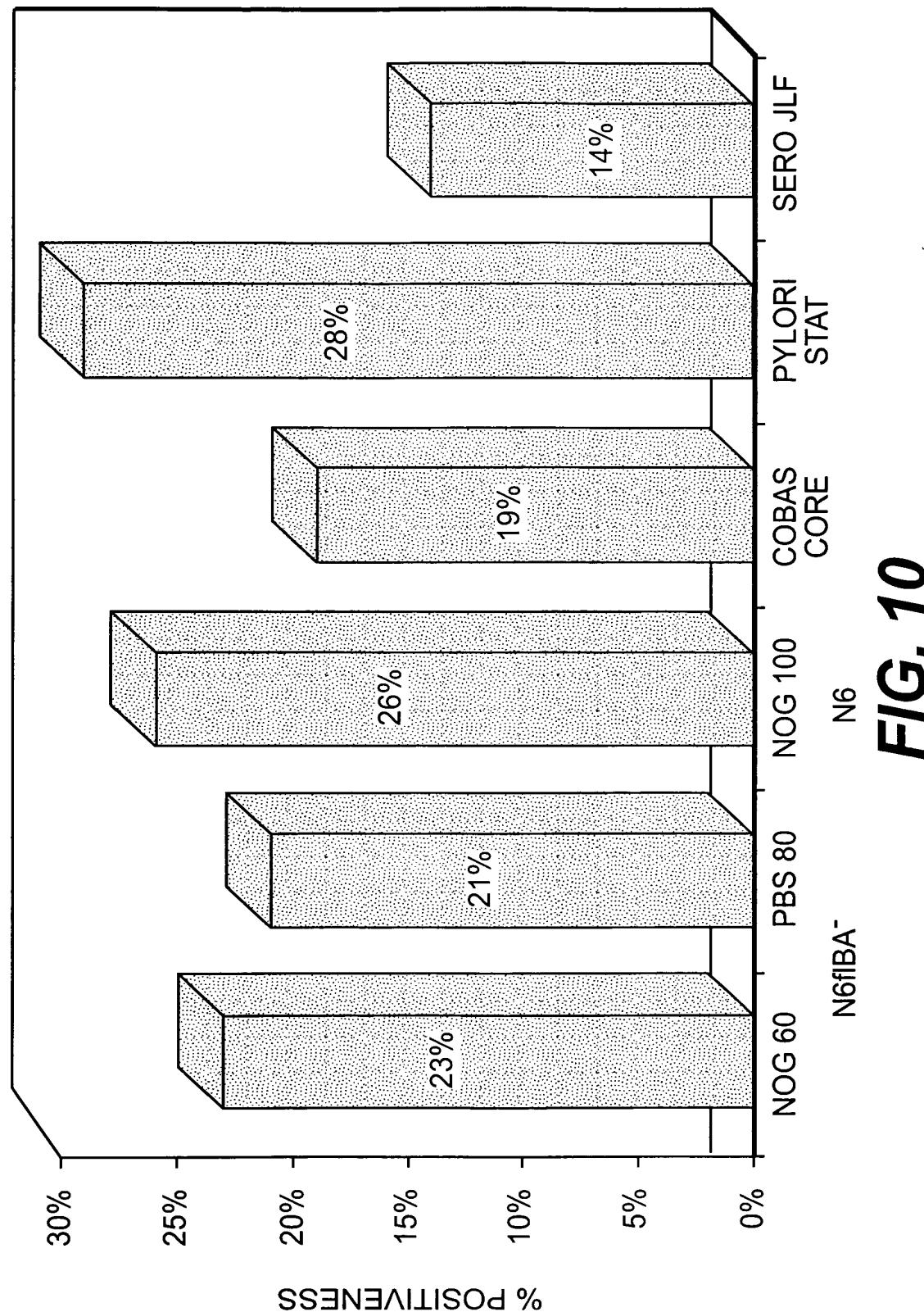
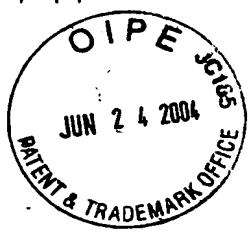
**FIG. 8**

# Replacement Sheet



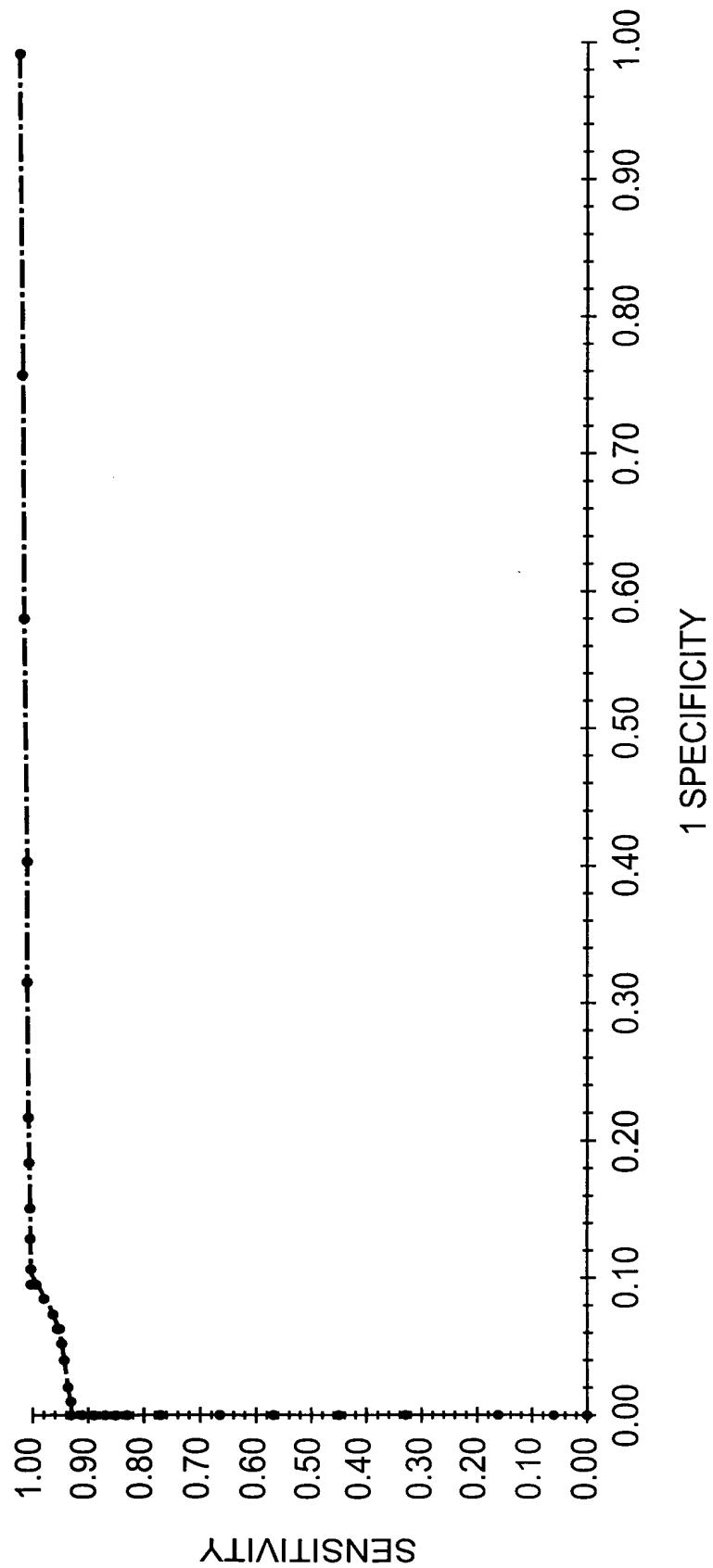
**FIG. 9**

Replacement Sheet



**FIG. 10**

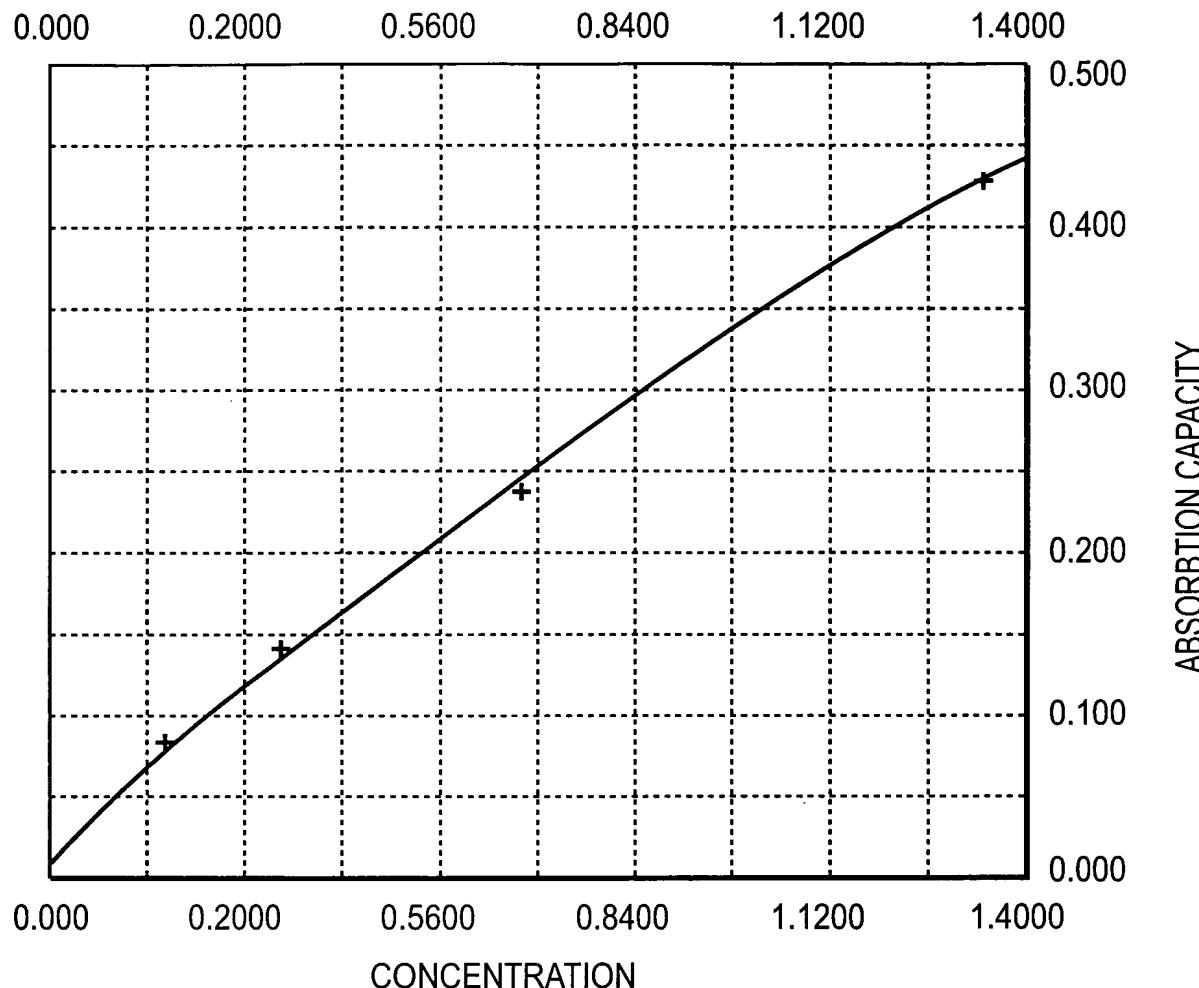
Replacement Sheet



**FIG. 11**



## Replacement Sheet

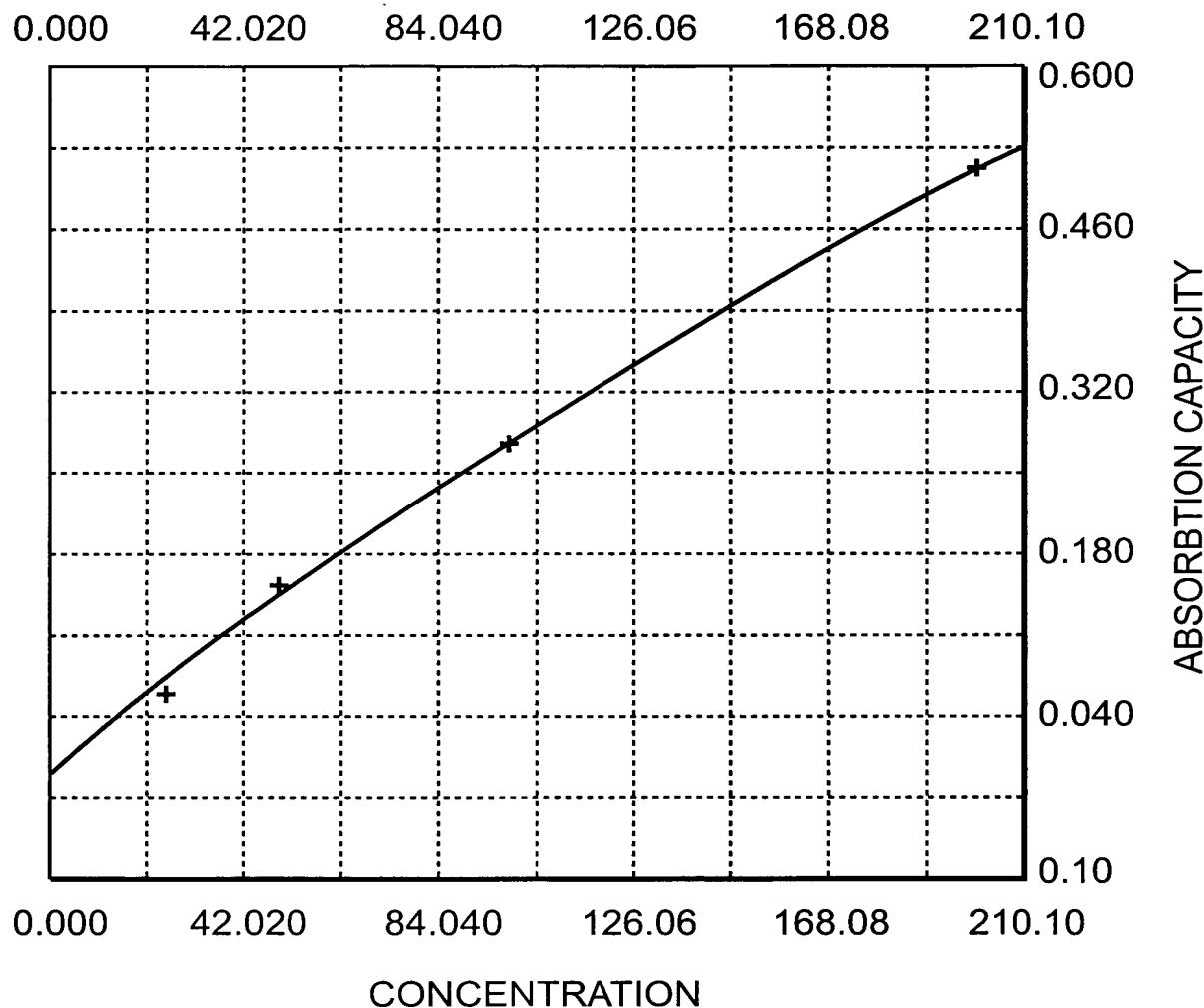


TYPE OF EXTRACT	DO @ 750nm	CONCENTRATION IN mg/ml
<u>GLYCINE</u> (AFTER CENTRIFUGATION FOR 15 min @ 3000 g)	0.028	0.284
<u>N-OCTYL-GLYCOSIDE</u>	0.087	1.004
<u>SUPERNATANT 1</u> (AFTER 1st PBS WASHING)	0.059	0.844
<u>SUPERNATANT 2</u> (AFTER 2nd PBS WASHING)	0.015	0.1105

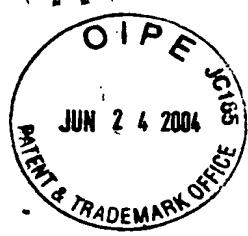
**FIG. 12**



## Replacement Sheet



**FIG. 13(A)**



## Replacement Sheet

TYPE OF EXTRACT	DO @ 760 nm	CONCENTRATION IN µg/ml
<u>GLYCINE</u> (AFTER CENTRIFUGATION FOR 15min @ 3000g)	0.279	202.86
<u>N-OCTYL-GLUCOSIDE</u>	0.243	873.99
<u>SUPERNATANT 1</u> (AFTER 1ST PBS WASHING)	0.361	539.2
<u>SUPERNATANT 2</u> (AFTER 2ND PBS WASHING)	0.218	77.875

TYPE OF EXTRACT	DO @ 760 nm	CONCENTRATION IN µg/ml
<u>GLYCINE RESIDUE</u> (AFTER 15 min OF CENTRIFUGATION AT 3000g)	0.099	297.5
<u>GLYCINE RESIDUE</u> (AFTER EXTRACTION)	0.093	2778.7
<u>N-OCTYL-GLUCOSIDE</u> (AFTER EXTRACTION)	0.275	972.0

**FIG. 13B**